



RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/089,508

DATE: 04/17/2002 TIME: 11:26:53

Input Set : A:\seqlist.txt

ENTERED Output Set: N:\CRF3\04172002\J089508.raw 4 <110> APPLICANT: Abuin, Alejandro

```
Clapham, Hohn
      8 <120> TITLE OF INVENTION: Transgenic Rodent Comprising A Polynucleotide Encoding A
             Human UCP3 Polypeptide
     12 <130> FILE REFERENCE: P32426
C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/089,508
C--> 14 <141> CURRENT FILING DATE: 2002-03-28
     14 <160> NUMBER OF SEO ID NOS: 2
     16 <170> SOFTWARE: FastSEQ for Windows Version 3.0
     18 <210> SEQ ID NO: 1
     19 <211> LENGTH: 936
    20 <212> TYPE: DNA
    21 <213> ORGANISM: Homo sapiens
    23 <400> SEQUENCE: 1
    24 atggttggac tgaagcette agacgtgeet eccaecatgg etgtgaagtt eetgggggea
                                                                                60
    25 ggcacagcag cctgttttgc tgacctcgtt acctttccac tggacacagc caaggtccgc
                                                                               120
    26 ctgcagatcc agggggagaa ccaggcggtc cagacggccc ggctcgtgca gtaccgtggc
                                                                               180
        gtgctgggca ccatcctgac catggtgcgg actgagggtc cctgcagccc ctacaatqqq
                                                                               240
    28 ctggtggccg gcctgcagcg ccagatgagc ttcgcctcca tccgcatcgg cctctacgac
                                                                               300
    29 tecgteaage aggtgtacae ceceaaagge geggacaaet ceageeteae taeeeggatt
                                                                               360
    30 ttggccggct gcaccacagg agccatggcg gtgacctgtg cccagcccac agatgtggtg
                                                                               420
    31 aaggteegat tteaggeeag catacacete gggeeateea ggagegaeag aaaatacage
                                                                               480
    32 gggactatgg acgcctacag aaccategee agggaggaag gagteagggg cetgtggaaa
                                                                               540
    33 ggaactttgc ccaacatcat gaggaatgct atcgtcaact gtgctgaggt ggtgacctac
                                                                               600
    34 gacatectea aggagaaget getggaetae eacetgetea etgacaaett eecetgeeae
                                                                               660
    35 tttgtctctg cctttggage cggcttctgt gccacagtgg tggcctcccc ggtggacgtg
    36 gtgaagaccc ggtatatgaa ctcacctcca ggccagtact tcagccccct cgactgtatg
    37 ataaagatgg tggcccagga gggccccaca gccttctaca agggatttac accctccttt
                                                                              840
    38 ttgcgtttgg gatcctggaa cgtggtgatg ttcgtaacct atgagcagct gaaacgggcc
                                                                              900
    39 ctgatgaaag tocagatgtt acgggaatca ccgttt
                                                                              936
    41 <210> SEQ ID NO: 2
    42 <211> LENGTH: 312
    43 <212> TYPE: PRT
    44 <213> ORGANISM: Homo sapiens
    46 <400> SEQUENCE: 2
    47 Met Val Gly Leu Lys Pro Ser Asp Val Pro Pro Thr Met Ala Val Lys
    48
                                            10
    49
        Phe Leu Gly Ala Gly Thr Ala Ala Cys Phe Ala Asp Leu Val Thr Phe
        Pro Leu Asp Thr Ala Lys Val Arg Leu Gln Ile Gln Gly Glu Asn Gln
    52
                                   40
        Ala Val Gln Thr Ala Arg Leu Val Gln Tyr Arg Gly Val Leu Gly Thr
```

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```
55 Ile Leu Thr Met Val Arg Thr Glu Gly Pro Cys Ser Pro Tyr Asn Gly
                     7.0
                                       75
57 Leu Val Ala Gly Leu Gln Arg Gln Met Ser Phe Ala Ser Ile Arg Ile
58
                 85
                                    90
59 Gly Leu Tyr Asp Ser Val Lys Gln Val Tyr Thr Pro Lys Gly Ala Asp
60
       100
                                105
61 Asn Ser Ser Leu Thr Thr Arg Ile Leu Ala Gly Cys Thr Thr Gly Ala
                            120
   Met Ala Val Thr Cys Ala Gln Pro Thr Asp Val Val Lys Val Arg Phe
                         135
                                          140
65 Gln Ala Ser Ile His Leu Gly Pro Ser Arg Ser Asp Arg Lys Tyr Ser
                     150
                                        155
   Gly Thr Met Asp Ala Tyr Arg Thr Ile Ala Arg Glu Glu Gly Val Arg
68
                 165
                                    170
69 Gly Leu Trp Lys Gly Thr Leu Pro Asn Ile Met Arg Asn Ala Ile Val
70
              180
                                185
71 Asn Cys Ala Glu Val Val Thr Tyr Asp Ile Leu Lys Glu Lys Leu Leu
72
          195
                             200
                                               205
73
   Asp Tyr His Leu Leu Thr Asp Asn Phe Pro Cys His Phe Val Ser Ala
74
      210
                         215
75 Phe Gly Ala Gly Phe Cys Ala Thr Val Val Ala Ser Pro Val Asp Val
76 225
                     230
   Val Lys Thr Arg Tyr Met Asn Ser Pro Pro Gly Gln Tyr Phe Ser Pro
77
                  245
78
                                    250
79 Leu Asp Cys Met Ile Lys Met Val Ala Gln Glu Gly Pro Thr Ala Phe
80
              260
                                265
   Tyr Lys Gly Phe Thr Pro Ser Phe Leu Arg Leu Gly Ser Trp Asn Val
81
                            280
                                              285
   Val Met Phe Val Thr Tyr Glu Gln Leu Lys Arg Ala Leu Met Lys Val
                         295
85 Gln Met Leu Arg Glu Ser Pro Phe
                     310
```



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VERIFICATION SUMMARY
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L:14 M:270 C: Current Application Number differs, Replaced Current Application No L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date